

## AMENDMENTS TO THE SPECIFICATION

Please amend the Brief Description of the Drawings for Figure 1L and 1M on pages 6 and 7, respectively as follows:

**Figure 1L:** The complete predicted amino acid sequence of VCIP (SEQ ID No. 13) is shown. The putative transmembrane segments are underlined. The N-linked glycosylation site is indicated by an asterisk. The lipid phosphatase catalytic domain is indicated by bold letters and underlined and the RGD motif is boxed.

**Figure 1M:** The amino acid sequence of human VCIP (top; SEQ ID No. 14) is aligned with other proteins containing lipid phosphatase-like catalytic motifs (SEQ ID Nos. 15-19).

Please amend the Brief Description of the Drawings for Figure 14A-H on page 13 as follows:

**Figures 14A-H** show schematic diagrams of various pLNCX2 retroviral constructs used in Figures 14-19. Figure 14H shows the amino acid sequence of PAP2b (SEQ ID No. 42) in the construct The relative positions of RGD, RGE, and RAD within the constructs are indicated. Arrows indicate the direction of transcription (5'-3'). Asterisk indicates the location of lipid phosphatase-dead motif.

## **AMENDMENT TO THE SEQUENCE LISTING**

Please replace the Sequence Listing mailed on August 18, 2005  
with the enclosed Sequence Listing